#11

1600

1632

P.6

RAW SEQUENCE LISTING
PATENT APPLICATION: **US/09/771,208A**

DATE: 11/21/2002
TIME: 10:06:45

Input Set : **A:\407T-923710US.ST25.txt**
Output Set: **N:\CRF4\11212002\I771208A.raw**

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 3 <110> APPLICANT: MEDRANO, JUAN
 4        BRADFORD, ERIC
 5        HORVAT, SIMON
 7 <120> TITLE OF INVENTION: CLONING OF A HIGH-GROWTH GENE
 9 <130> FILE REFERENCE: 407T-923710US
11 <140> CURRENT APPLICATION NUMBER: US 09/771,208A
12 <141> CURRENT FILING DATE: 2001-01-26
14 <150> PRIOR APPLICATION NUMBER: US 08/999,477
15 <151> PRIOR FILING DATE: 1997-12-29
17 <160> NUMBER OF SEQ ID NOS: 21
19 <170> SOFTWARE: PatentIn version 3.0
21 <210> SEQ ID NO: 1
22 <211> LENGTH: 1667
23 <212> TYPE: DNA
24 <213> ORGANISM: Mus musculus
26 <400> SEQUENCE: 1
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31 ggaagccaga gacaagcagg tactccgctc cctgcgtctg gagctgggtg ccgaggtact   180
33 ggtggaagga ctggttcttc agtaccttta ccaggaagga attttgacag aaaaccacat   240
35 tcaagaaatc aaagctcaaa ccacaggcct ccggaagaca atgctgttgc tggacatcct   300
37 gccttccagg ggccccaaag cttttgacac cttcctcgat tccctccagg aatttccctg   360
39 ggtaagagag aagctggaga aggcgagaga ggaagtctca gccgagctgc ctacaggtga   420
41 ctggatggcc ggaatcccct cacacatcct cagcagctcg ccatcagacc agcagattaa   480
43 ccagctggct cagaggctag gcccggagtg ggagcccgtg gtcctgtctc tgggactgtc   540
45 ccagaccgac atctaccgct gcaaggccaa ccatccccac aacgtgcatt cgcaggtggt   600
47 ggaggccttt gtccgctggc gccagcgttt tgggaagcag gccaccttcc taagcttaca   660
49 caagggcctc caggcaatgg aggctgatcc ctccctgctc cagcacatgc tggagtgacc   720
51 tgacccccccc ccgcgcccccc ccccacttg ctgtgggggt ggtggggcgt gggttcccaa   780
53 gtcacactgg ctgaaccgga cttttctcag caggtggctt tgttctgggc ttttcagtga   840
55 tctgtttacg gaaagagatc gtccaccact cactcaacca tcgattggct ttaattgctt   900
57 gaagactgcg ctgttgtaac tatggtttgg aactttgtgg ctggccttta acaggaggcc   960
59 agaaaaaaca caacacccac cctacccaac cccccaaaaa atcatgctac agcatcgaat  1020
61 gcaggtgtcc tgcatacaag gcagctacac ttgtgttgcc tggagactgg attgtgcatt  1080
63 tagctcttca taatggtgat gataataaaa aagcaaattg tgatatagaa tgtgcctctt  1140
65 tcaatgagag agtattatat cacacacaca cacacacaca cacacacaca tacacacaca  1200
67 cacaccaatc ttctgttgca tagacggagg gtgtaaaaat atgggagtgg agcaagattg  1260
69 atagcagtca tgtgacgacg gagataaata actcaggcag gatgtataga ttaagcatga  1320
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73 gtatgaggag tctctgtcca tttgttaaag gcattggatc agagacaaga gggctcagtg  1440
75 tttctcttga ggcctgaatg gctgaaggcg gtgagttccc gaggggcgtc atgggttgtc  1500
77 cagcctttca ttaactgcac atagtgttag ccagacaggt gtacgtgttt gtcatcccat  1560
79 ctaagagact gaagcaggag gatcacctgt acatgactgc ttctttcaac attttaaaat  1620
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RAW SEQUENCE LISTING   DATE: 11/21/2002
PATENT APPLICATION: **US/09/771,208A**   TIME: 10:06:45

Input Set : **A:\407T-923710US.ST25.txt**
Output Set: **N:\CRF4\11212002\I771208A.raw**

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87  <213> ORGANISM: Mus musculus
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94  aaccacattc aacaaatcaa agctcaaacc acaggcctcc ggaagacaat gctgttgctg    180
96  gacatcctgc cttccagggg ccccaaagct tttgacacct tcctcgattc cctccaggaa    240
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100 acag                                                                304
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104 <211> LENGTH: 160
105 <212> TYPE: DNA
106 <213> ORGANISM: bovine sp
108 <400> SEQUENCE: 3
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111 ggtactggtg gagggctag tcctccagta tctttatcag gaagggtct tgacagaaag     120
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117 <211> LENGTH: 539
118 <212> TYPE: PRT
119 <213> ORGANISM: Mus musculus
121 <400> SEQUENCE: 4
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126 Ser Gly Leu Asn Val Ala Leu Leu Glu Ala Arg Gly Ser Glu Arg Leu
127             20                  25                  30
129 Glu Ala Arg Gly Leu Glu Gly Leu Leu Glu Gly Leu Tyr Ala Leu Ala
130         35                  40                  45
132 Gly Leu Val Ala Leu Leu Glu Val Ala Leu Gly Leu Gly Leu Tyr Leu
133     50                  55                  60
135 Glu Val Ala Leu Leu Glu Gly Leu Asn Thr Tyr Arg Leu Glu Thr Tyr
136 65                  70                  75                  80
138 Arg Gly Leu Asn Gly Leu Gly Leu Tyr Ile Leu Glu Leu Glu Thr His
139             85                  90                  95
141 Arg Gly Leu Ala Ser Asn His Ile Ser Ile Leu Glu Gly Leu Asn Gly
142             100                 105                 110
144 Leu Ile Leu Glu Leu Tyr Ser Ala Leu Ala Gly Leu Asn Thr His Arg
145         115                 120                 125
147 Thr His Arg Gly Leu Tyr Leu Glu Ala Arg Gly Leu Tyr Ser Thr His
148     130                 135                 140
150 Arg Met Glu Thr Leu Glu Leu Glu Leu Glu Ala Ser Pro Ile Leu Glu
151 145                 150                 155                 160
153 Leu Glu Pro Arg Ser Glu Arg Ala Arg Gly Gly Leu Tyr Pro Arg Leu
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156 Tyr Ser Ala Leu Ala Pro His Glu Ala Ser Pro Thr His Arg Pro His
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159 Glu Leu Glu Ala Ser Pro Ser Glu Arg Leu Glu Gly Leu Asn Gly Leu
160         195             200             205
162 Pro His Glu Pro Arg Thr Arg Pro Val Ala Leu Ala Arg Gly Gly Leu
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165 Leu Tyr Ser Leu Glu Gly Leu Leu Tyr Ser Ala Leu Ala Ala Arg Gly
166 225             230             235             240
168 Gly Leu Gly Leu Val Ala Leu Ser Glu Arg Ala Leu Ala Gly Leu Leu
169         245             250             255
171 Glu Pro Arg Thr His Arg Gly Leu Tyr Ala Ser Pro Thr Arg Pro Met
172         260             265             270
174 Glu Thr Ala Leu Ala Gly Leu Tyr Ile Leu Glu Pro Arg Ser Glu Arg
175         275             280             285
177 His Ile Ser Ile Leu Glu Leu Glu Ser Glu Arg Ser Glu Arg Ser Glu
178         290             295             300
180 Arg Pro Arg Ser Glu Arg Ala Ser Pro Gly Leu Asn Gly Leu Asn Ile
181 305             310             315             320
183 Leu Glu Ala Ser Asn Gly Leu Asn Leu Glu Ala Leu Ala Gly Leu Asn
184         325             330             335
186 Ala Arg Gly Leu Glu Gly Leu Tyr Pro Arg Gly Leu Thr Arg Pro Gly
187         340             345             350
189 Leu Pro Arg Val Ala Leu Val Ala Leu Leu Glu Ser Glu Arg Leu Glu
190         355             360             365
192 Gly Leu Tyr Leu Glu Ser Glu Arg Gly Leu Asn Thr His Arg Ala Ser
193         370             375             380
195 Pro Ile Leu Glu Thr Tyr Arg Ala Arg Gly Cys Tyr Ser Leu Tyr Ser
196 385             390             395             400
198 Ala Leu Ala Ala Ser Asn His Ile Ser Pro Arg His Ile Ser Ala Ser
199         405             410             415
201 Asn Val Ala Leu His Ile Ser Ser Glu Arg Gly Leu Asn Val Ala Leu
202         420             425             430
204 Val Ala Leu Gly Leu Ala Leu Ala Pro His Glu Val Ala Leu Ala Arg
205         435             440             445
207 Gly Thr Arg Pro Ala Arg Gly Gly Leu Asn Ala Arg Gly Pro His Glu
208         450             455             460
210 Gly Leu Tyr Leu Tyr Ser Gly Leu Asn Ala Leu Ala Thr His Arg Pro
211 465             470             475             480
213 His Glu Leu Glu Ser Glu Arg Leu Glu His Ile Ser Leu Tyr Ser Gly
214         485             490             495
216 Leu Tyr Leu Glu Gly Leu Asn Ala Leu Ala Met Glu Thr Gly Leu Ala
217         500             505             510
219 Leu Ala Ala Ser Pro Pro Arg Ser Glu Arg Leu Glu Leu Glu Gly Leu
220         515             520             525
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227 <212> TYPE: DNA
228 <213> ORGANISM: Artificial
230 <220> FEATURE:
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239 <212> TYPE: DNA
240 <213> ORGANISM: Artificial
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243 <223> OTHER INFORMATION: PCR primer
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251 <212> TYPE: DNA
252 <213> ORGANISM: Artificial
254 <220> FEATURE:
255 <223> OTHER INFORMATION: PCR primer
257 <400> SEQUENCE: 7
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262 <211> LENGTH: 21
263 <212> TYPE: DNA
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266 <220> FEATURE:
267 <223> OTHER INFORMATION: PCR primer
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276 <213> ORGANISM: Mus musculus
278 <400> SEQUENCE: 9
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283 cttcttagct ggtagcactc acttttgaag gctgtaaagc cttcaatctg tgggtcctac  180
285 ccctttggca aaccttgatc tccaaagtta cataagcaca ggcacacact tccacttcct  240
287 ctgaggtttt ctaccaagaa aggatcaacc attcataaaa tgttggtcct agtgaaccct  300
289 gcacattgta gaggcttaaa aagtttaatt tgggcctcca actcactaca caggaactcc  360
291 agcgggatcc gcctgtccgt tcatgctaac cttcaccga catcttgttt ttaagtttac  420
293 agaaaacgtt agggacctaa agaaggtcat tacattacag tacattacag tacaacagaa  480
295 gttacaaagt agcaatgagg ggcttgggga tttagctcag tgctagagcg cttgcctagc  540
297 aagtgcaaga ccctaggttc ggtcctcagc tctgaaaaat caaaacaaaa caaaacaaag  600
299 tagcaatgat aataatttta tggttgaggg gtcaccatga tatgaggaac tgtattaaac  660
301 ggtcgctgca ttagggagga tgaggaccac tgtggggctc agctgaagga agtgagttgc  720
303 tggtgtaggg caccggagtg ctagatgtaa accggtttcc tgtctccctt ctaaggctga  780
305 ctgcaccact aattcctgcc tcccgtggag ggtgctttcc aggctccaag ccttcctgcc  840
307 atgttggaat gtgtcctgtg aaccatgaac cgagatcaat cttcctccc ttccatcacc   900
309 tctgccaggt ggtttggtca tagtactcag tagagtaagg aggctggaag atttactaca  960
311 cctgacaaag aaaaattaat ctgtatgatc tcaaaaaaaa aaaaaaaaa aacaccacca  1020
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RAW SEQUENCE LISTING   DATE: 11/21/2002
PATENT APPLICATION: **US/09/771,208A**   TIME: 10:06:45

Input Set : **A:\407T-923710US.ST25.txt**
Output Set: **N:\CRF4\11212002\I771208A.raw**

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315 acacttccac ttcctctgag gttttctacc aagaaaggat caaccattca taaaacgttg 1140
317 gtcttagtta tccctgcaca ttgtagaggc ttaaaaagtt aacttgggc ctccaactca 1200
319 ctacacagaa ctccagaggg atccgcctgt ccgttcatgc taacctttca ccgacatctt 1260
321 gtttttaagt ttacagaaaa cgttagggac ctaaagaagg taagcatcct gctaagttac 1320
323 tccctggctt tacacaggct ttctaaaact tgagtaagag gcatccttcc catcaaagat 1380
325 tccaggaaaa cagcctcccc cctccgcggc cacacatacg aatctatcgc tgacaaagcc 1440
327 cctgtaagct ggcttatgtc ctccctcgc gttcaccatt ctgtaagtgc atagaattat 1500
329 ttaagaggaa aaaaattact gtggataaaa attggttcgg ggccttggaa ttggccggtc 1560
331 gtgttgtgtt tccttccagg gccggcaggc ggggcaccag gcaaggcttg gaagccgcgc 1620
333 ctctctcaac ctctcctggc cacccttgcc caacttcccc atagacacag cttcaactaa 1680
335 aagtggccat tgacctttca agcttttgag cagtgggca acagaacagt atttcaaaga 1740
337 aaaatggtta tcgaaatttc gaatccggtt ttcccatgag tgtttttttt ttgtttgtt 1800
339 tttctgttaa aaaaaaaaa agtaggtcac attcaaagtg ggtcacgttt caggagccgg 1860
341 cgtgcctgga tgccggcgcg gaggctaggt ggcctcttac agagtgggag gtgagggtcc 1920
343 caataggaaa gaagtactgg gatcaatacg aactccgggt ccctggcttt gcaaggattc 1980
345 acagagacaa acgcaccagg cctgtgaccc cgcaccccac ccgggcacag gtaagggcac 2040
347 ctcctctgta gggtggccag ggtgggtctc ccgaagggca agcaggagtt gagctgagga 2100
349 ggaaaggaga agctgggcaa ggctgatgca ggggactacc agttggagct ccaggggga 2160
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367 accagcacca gcacccgcac cggctctctc cacccggctc ccttgaagcc tgcgcattag 2700
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371 aagttcctgg aaagcggcct ccgccgcggc cgggcggggc gcgaggggcg ggaggcgggg 2820
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393 cagggtgtc agcctggggc ggagagggg ggggaagct aggcgacgat ccctgggatt 3480
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407 cgccgctgag ccccccatcc tcgtctggag cgtgctccag gaagcggcag gagtgggggt 3900
409 gaggccgcct ccgaggcagg gatgcagcgg ctggcgcgcc gctagcgcac cgcagcaccc 3960
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 11/21/2002
PATENT APPLICATION: **US/09/771,208A**   TIME: 10:06:46

Input Set : **A:\407T-923710US.ST25.txt**
Output Set: **N:\CRF4\11212002\I771208A.raw**

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

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Seq#:20; N Pos. 132692,132693,132694,132695,132696,132697,132698,132699
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Invalid <213> Response:
Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

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Seq#:5,6,7,8,10,11,12,13,14,15,16,17,18,19,21
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